

FIG. 1

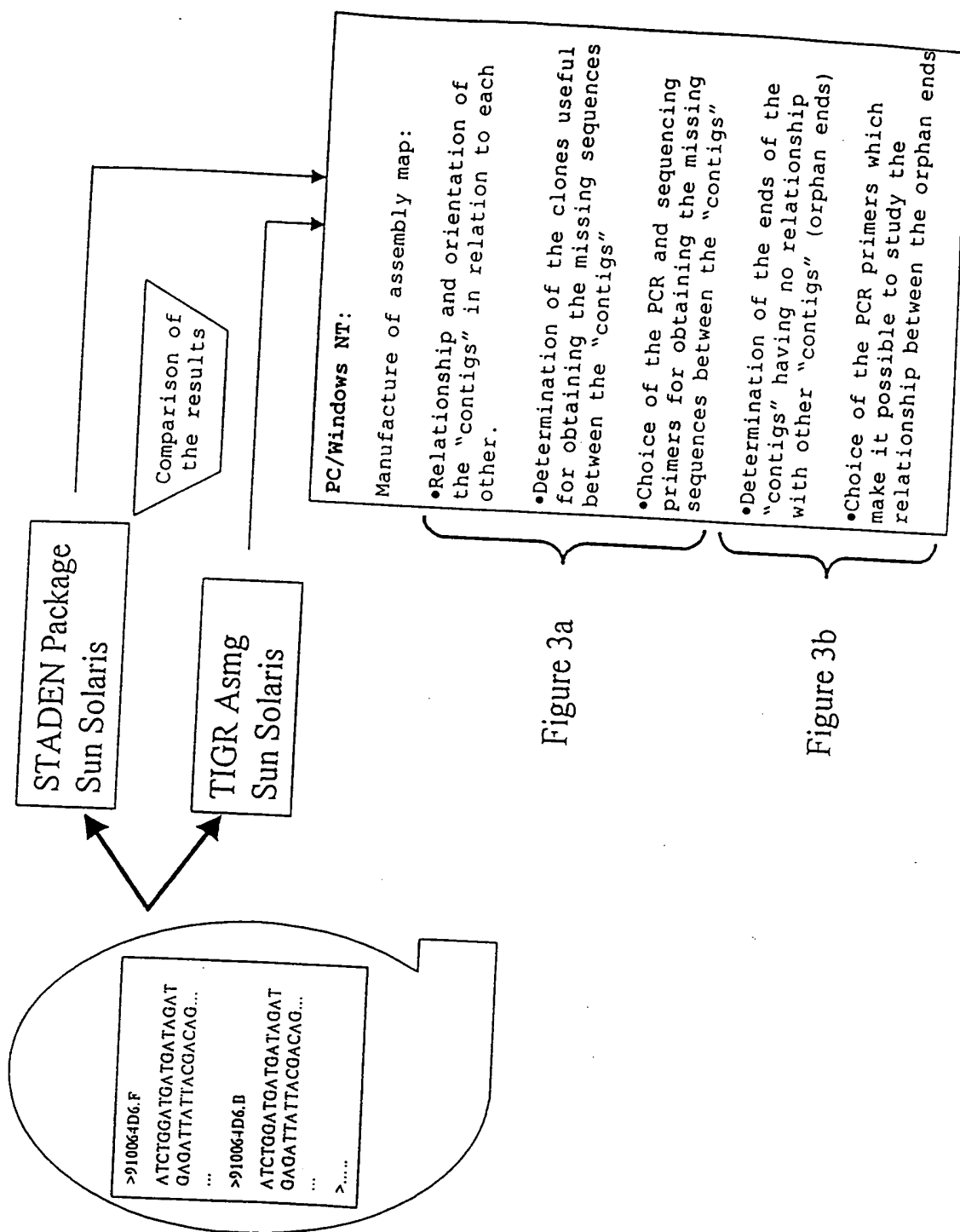
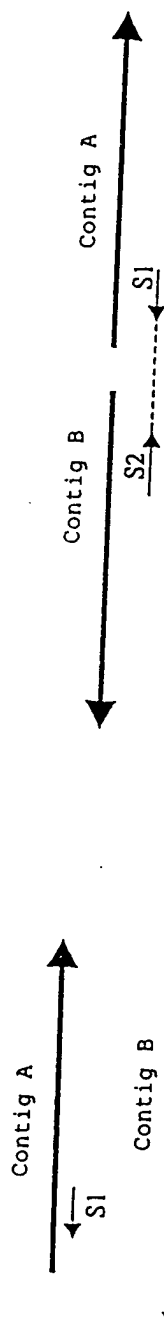


Figure 3a

Figure 3b

FIG. 2

a.

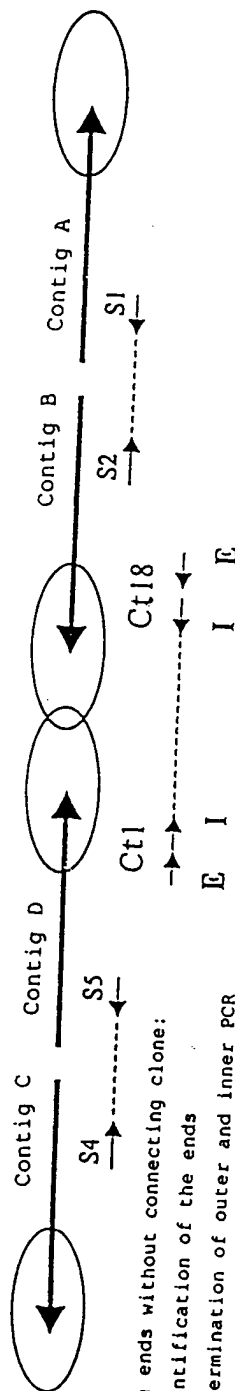


Distribution and position of the sequences on the contigs:  
 1- Define the arrangement of the contigs in relation to each other  
 2- Define the PCR primers which make it possible to fill the sequence

Statistical determination of the sequences:

- 1- Belonging to the same clone
- 2- Situated on two different contigs

b.



Contig ends without connecting clone:

- 1- Identification of the ends
- 2- Determination of outer and inner PCR primers for studying the relationships between the contigs

E: outer primers  
 I: inner primers

FIG. 3